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## AMENDMENTS TO THE CLAIMS

1-21 (Cancelled)

:

- 22. (Previously presented) An isolated polypeptide having at least 80% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, wherein said isolated polypeptide bas the ability to induce chondrocyte redifferentiation;
  - (b) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, lacking its associated signal peptide, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation; or
  - (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203581, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation.
- 23. (Previously presented) The isolated polypoptide of Claim 22 having at least 85% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation;
  - (b) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, lacking its associated signal peptide, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation; or
  - (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203581, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation.
- 24. (Previously presented) The isolated polypeptide of Claim 22 having at least 90% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation;

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- (b) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, lacking its associated signal peptide, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation; or
- (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203581, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation.
- 25. (Previously presented) The isolated polypeptide of Claim 22 having at least 95% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation;
  - (b) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, lacking its associated signal peptide, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation; or
  - (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203581, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation.
- 26. (Previously presented) The isolated polypeptide of Claim 22 having at least 99% amino acid sequence identity to:
  - (a) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation;
  - (b) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, lacking its associated signal peptide, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation; or
  - (c) the amino acid sequence of the rolypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203581, wherein said isolated polypeptide has the ability to induce chondrocyte redifferentiation.

27-32 (Cancelled)

33. (Previously presented) A chimeric polypeptide comprising a polypeptide according to Claim 22 fused to a heterologous polypeptide.

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- 34. (Previously presented) The chimeric polypeptide of Claim 33, wherein said heterologous polypeptide is an epitope tag or an Fc region of an immunoglobulin.
  - 35. (Previously presented) An isolated polypertide comprising:
  - (a) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2;
  - (b) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, lacking its associated signal peptide; or
  - (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203581.
- 36. (Previously presented) The isolated polypeptide of Claim 35 comprising the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2.
- 37. (Previously presented) The isolated polypeptide of Claim 35 comprising the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, lacking its associated signal peptide.
- 38. (Previously presented) An isolated polypeptide comprising a fragment of a polypeptide having the sequence of SEQ ID NO:2, wherein said fragment comprises amino acids 137-167 of SEQ ID NO:2.
- 39. (Previously presented) The isolated polypeptide of Claim 38, wherein said fragment consists essentially of amino acids 137-167 of SEQ ID NO:2.
- 40. (Previously presented) An isolated polypeptide comprising a fragment of a polypeptide having the sequence of SEQ ID NO:2, wherein said fragment comprises amino acid sequences from SEQ ID NO:2 selected from the group consisting of amino acids 57-91, 60-94, 54-88, 81-114, 78-111, 63-96, 51-84, 45-78, 48-81, 33-65, 66-99, 42-75, 135-169, 202-221, and 235-244.
- 41. (New) An isolated polypeptide having at least 80% amino acid sequence identity to:
  - (a) the amino acid sequence of the pclypeptide having the sequence of SEQ ID NO:2;
  - (b) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, lacking its associated signal peptide; or

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(c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203581

wherein said isolated polypeptide comprises a C1q domain signature sequence or a C1q domain protein sequence.

- 42. (New) The isolated polypeptide of Claim 41, wherein said C1q domain signature comprises amino acids 137-167 of SEQ ID NO:2.
- 43. (New) The isolated polypeptide of Claim 41, wherein said C1q domain protein sequence comprises an amino acid sequence from SEQ ID NO:2 selected from the group consisting of amino acids 57-91, 60-94, 54-88, 81-114, 78-111, 63-96, 51-84, 45-78, 48-81, 33-66, 66-99, 42-75, 135-169, 202-221, and 235-244.
- 44. (New) The isolated polypeptide of Claim 41, wherein said isolated polypeptide further comprises a sequence having homology to a subunit of collagen alpha 1(x).
- 45. (New) An isolated polypeptide having at east 80% amino acid sequence identity to:
  - (a) the amino acid sequence of the pclypeptide having the sequence of SEQ ID NO:2;
  - (b) the amino acid sequence of the polypeptide having the sequence of SEQ ID NO:2, lacking its associated signal peptide; or
  - (c) the amino acid sequence of the polypeptide encoded by the full-length coding sequence of the cDNA deposited under ATCC accession number 203581

wherein said isolated polypeptide comprises a sequence having homology to a subunit of collagen alpha 1(x).

- 46. (New) An antibody or antibody fragment which specifically binds to a polypeptide according to any of Claims 22, 33, 35, 41 or 45.
  - 47. (New) A composition comprising:
    a peptide according to any of Claims 22, 33, 35, 41 or 45; and
    a pharmaceutically acceptable carrier.

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